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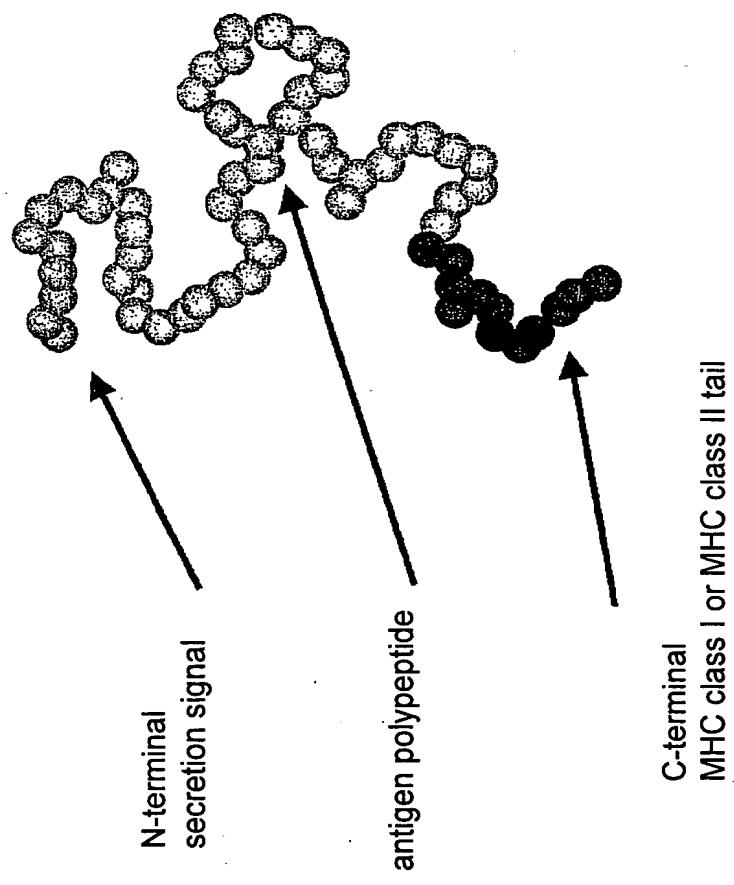
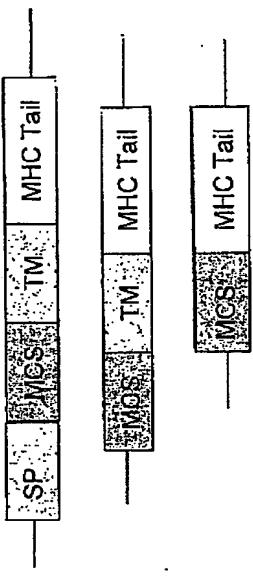


Fig. 1

**Fig. 2**

cassettes containing cloning sites (MCS) for expression of MHC fusion proteins of the invention



cassettes containing antigens cloned therein for expression of MHC fusion proteins of the invention

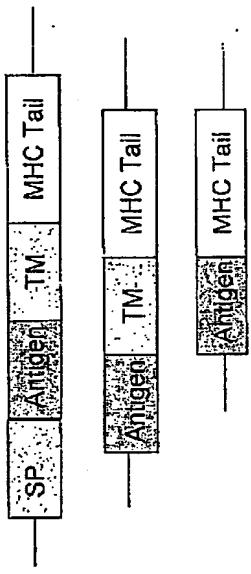
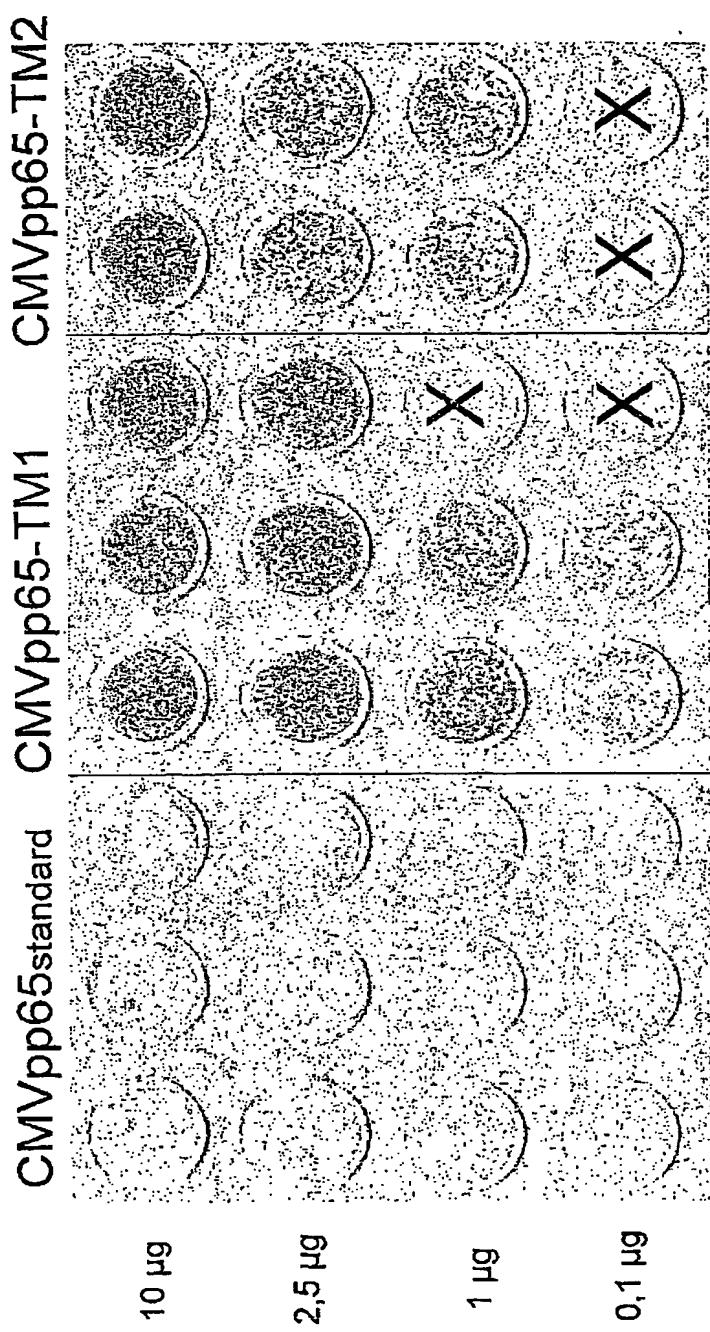
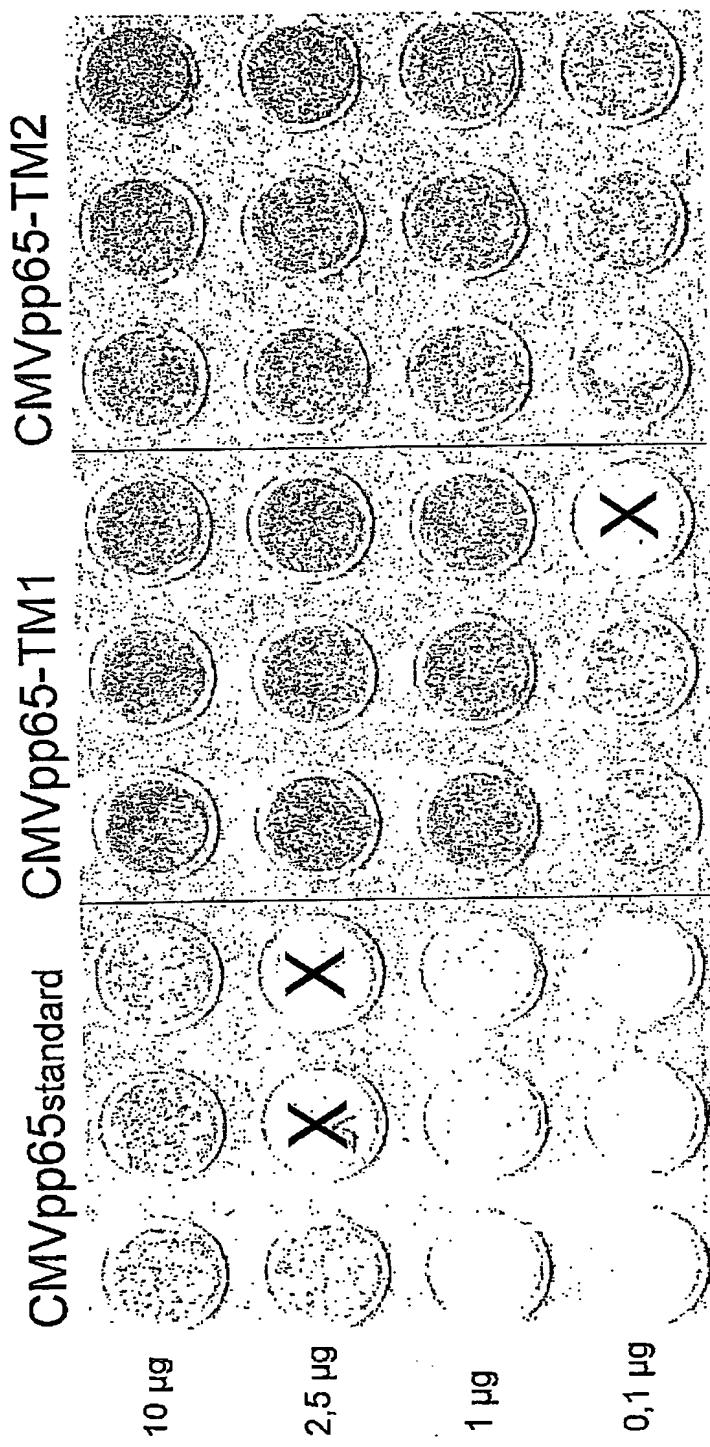


Fig. 3



**Fig. 4**

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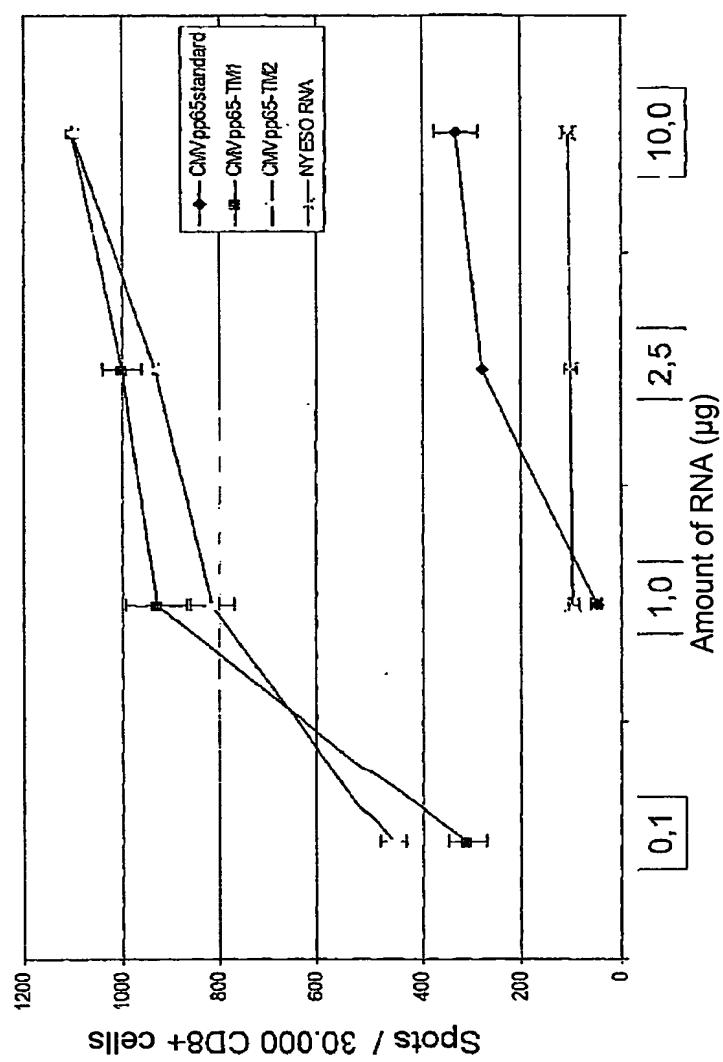
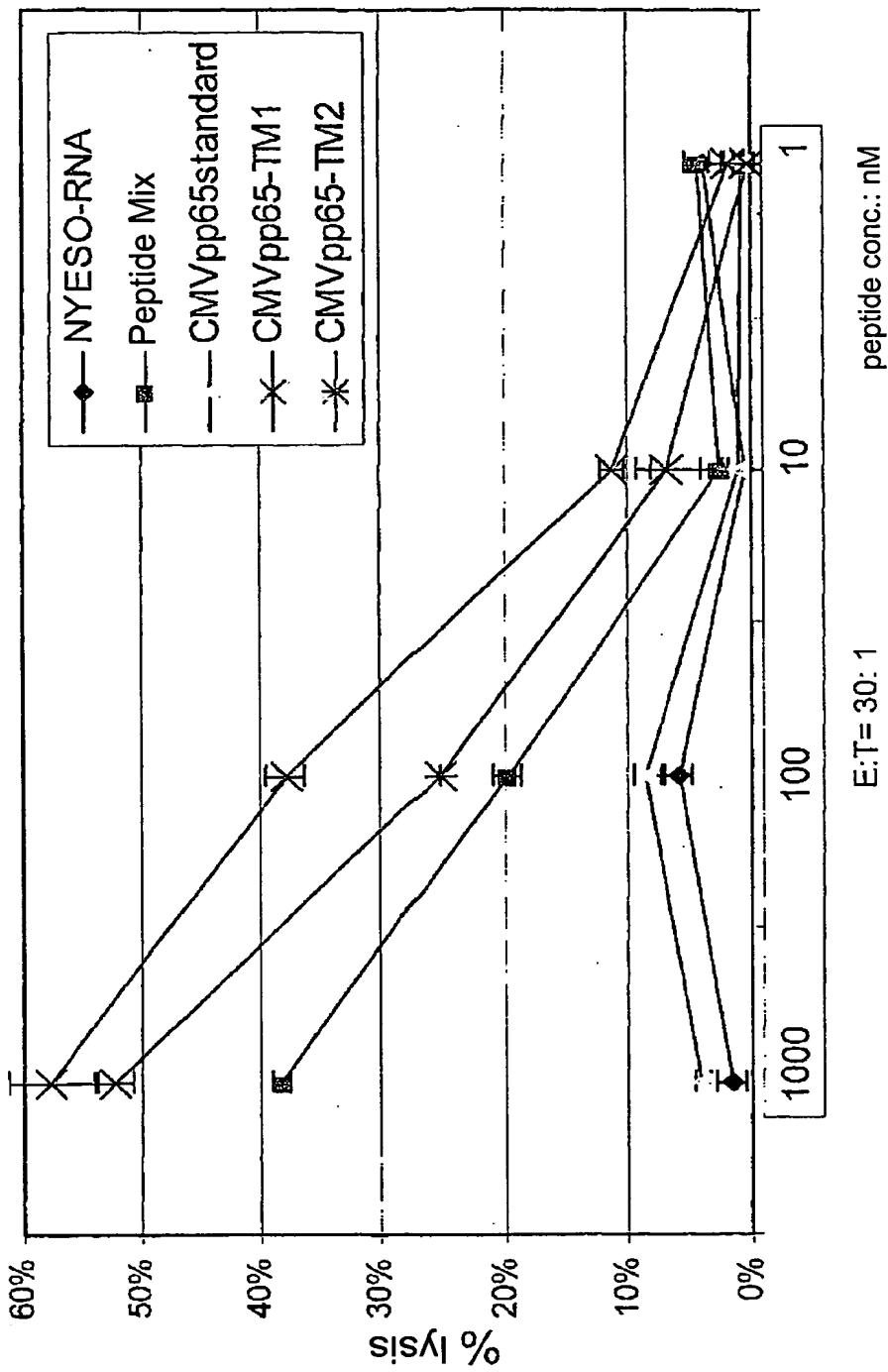


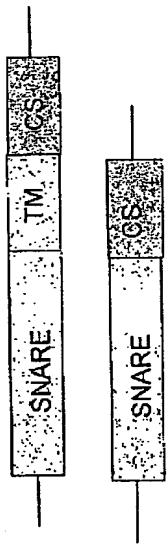
Fig. 5

Fig. 6

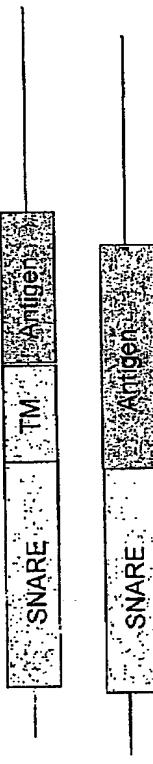


**Fig. 7**

cassettes containing cloning sites (CS) for expression of SNARE fusion proteins of the invention



cassettes containing antigens cloned therein for expression of SNARE fusion proteins of the invention



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			CSELENWTRATEKQVIGQYKVYLESCEDVPSCEKTLFEMHTVIGSDVEELDTYTRNPOEFMR PHIERNGTIVLCPKNMIIKPGKISHMLDVAFTSHEHFGILCPKSIPGSLISGNLLNMGQ ITLEVOA-TRETELVRDPEVAAL-FFDIDLLORGQYSEHPTFTSQRIQGRKTMXHTK DREDEGAQGGDDVVTSSESDDELVTERKTPTRVTCGAGASTSGRKRSASSATA QHLYQEFFFNWANDIVTPELEGGTQPAQPKGRURQDQALEFCIAST:TKKHRCGSQS RKLSSGVMTGRKIAESTVLPEDTDDESDNEINBANEVTNPWQAGLFLRNLYPMVATYQG LGLLFGAGLFLYFNRNQKHSGLQPRGFSL
64	PRAME variant 1	1 atggacgcaa ggcttgtt ggtttccatt cagagcugat acatcagcat gagtgtgtgg 61 accagccaa ggacactgt ggatggatggaa gggcgagcc -gctgtgaagg tgaggccctgg 121 ccatttgcgg cccttgagggt qctgcccaagg gagcttcggc cgttacactttt catggccggcc 181 ttgtacggaa gacacagcca gacccttggaa gcaatgtgc aggcctggcc cttcacctgc 241 ctccctccgg gagggtgtgt gatggatggaa cttccacca cttccacca tggagccctt cttttttttttt 301 ctgtatggac ttgtatgtct cttgtatgtct cttgtccca ggggtccca ggggtccca cttttttttttt 361 gtgtctggat tacggaaa ctctccatcg gacttggaa ctgtatggc tggaaatccatgg 421 gucaglctgt actcatttcc agggccggaa gggatggaa gggatggaa gggatggaa 481 tgatgtttt tgatgtttt tgatgtttt tgatgtttt tgatgtttt tgatgtttt 541 ctgttcctca aggaaatggc ctgtatgtttt tgatgtttt tgatgtttt tgatgtttt 601 cggaaaatggaa atgtatcag ctgtatcag ctgtatcag ctgtatcag ctgtatcag 661 caggatataca agatgtatcc gaaatggatctt cttttttttt cttttttttt 721 acttgtatcc ggaatgtttt cttttttttt cttttttttt 781. aatcttgctta gactctctctt ctccacacat catgcacatc catgcacatc 841 ggagggcaat atatgcacca ttccacatc ctttccatc ctttccatc 901 ctctatgtgg actctatgtttt ttctttatggaa gggatggatgg 961 atgaaacctt tggaaacctt cttttttttt 1021 catctglccc agatgtccgg cttttttttt 1081 ctgaccgtatg taatggccgg gcccccccaaa gttttttttt 1141 cttttttttt gttttttttt 1201 tccttgcgc actgtctccca gttttttttt 1261 tccttgcgc actgtctccca gttttttttt 1321 tttttttttt 1381 gcctatctgc atggcagct: cttttttttt 1441 tttttttttt 1501 atctgtgtgc cttttttttt 1 atgggtctccg acgtgtccggaa cttttttttt 61 ggccgggggt gtgccttgcgg tttttttttt 121 ggccgggggt ttatggccgg tttttttttt 181 cggcoaccyc cggccggccgg 241 cggcaqccgc acggaggagg 301 actccacacaat cccqdaqccctg 1 atgggtctccg acgtgtccggaa cttttttttt 61 ggccgggggt gtgccttgcgg tttttttttt 121 ggccgggggt ttatggccgg tttttttttt 181 cggcoaccyc cggccggccgg 241 cggcaqccgc acggaggagg 301 actccacacaat cccqdaqccctg	
65	WT1 variant C	1 atgggtctccg acgtgtccggaa cttttttttt 61 ggccgggggt gtgccttgcgg tttttttttt 121 ggccgggggt ttatggccgg tttttttttt 181 cggcoaccyc cggccggccgg 241 cggcaqccgc acggaggagg 301 actccacacaat cccqdaqccctg	



6  
Fig

Ser#	Seq ID	Type	Name	Sequence
No				
15	MHC	HLA-A		PSSOPTIPIVGLVLFIAVITGAVYAAVMWRRKSSDRKGGSYSQAASSDSAQGSDVSLTACKV
16	MHC	HLA-B		GSYSQAAASSDSQGSDVSLTACKV
17	MHC	HLA-B		PSSOPTIPIVGLVLAGLVAVVVGAVYAAVMCRRKSSGGKGSYSUAAACSDSAQGSDVSLITA
18	MHC			GSYSQAAACSDSQQGSDVSLITA
19	MHC	HLA-C		PSSOPTIPIVGLVAGLAVLAVLGAVMVAIVMCRKSQGRRGSCSQAISSNSAQGSDESPLITCKA
20	MHC			SAQGSDESPLITCKA
21	MHC	HLA-E		PSSOPTIPIVGLVLLGVSSEGVAVAVIWRKSSGCGGGSYSKAENSDSAQGSSESHSL
22	MHC			GSKSYKAENSDSAQGSSESHSL
23	MHC	HLA-F		QSPOPTIPIVGLVAVLVLVIGAVWTRGVVAAMWRRKSSDENRGYSQAAVTDQAQG3GV3LTANKV
24	MHC			BNRGSYSDAAVVDSAQGSGVSLVANKV
25	MHC	HLA-Dra		VVCALGLTVGVYGTITGTTFIIKGILRKSNAAERGPL
26	MHC			RKSNAEAERGPL
27	MHC	HLA-DRB		MLSGVGGFVQLGILTFAGLFLYFRNGKGHSGLQPRGFELIS
28	MHC			GHSGLQPRGFELIS
29	MHC	HLA-Dqa		WVOALGLSYGLMIVTVGTVLIIQGLRSVGASRHQGPL
30	MHC			VGNMRHQGPL
31	MHC	HLA-DCB		MLSGIGGGFVQLGILTFAGLFLYFRNGKGHSGLQPRGFELIS
32	MHC			RSGQGLIH
33	MHC	HLA-Dpa		VLCAALGLVGLVGVITVGTVLIKSLSDCHDPAQGTI
34	MHC			RSQHDPAQGTI
35	MHC	HLA-DBp		TITGAGGFVQLGILTCGGVGMMEERRSKVQRGSA
36	MHC			SKKVQRGSA
37	MHC	CD1a		FILIAVIVPLLLILGLALWERKRCFC
38	MHC			RKRCFC
39	MHC	CD1b		IYIAITIVSLLILCLIAWMMRRSYONIP
40	MHC			RRRSYONIP
41	MHC	CD1c		WIAAUUVVPLVLLVIVLWKKHCSYQDIL
42	MHC			KKHCSCYQDIL

Fig. 10

SEQ ID NO	Type	Name	Sequence
4.3	SNARE	Cis-golg1 SNARE p28	MAAGTSSYYWE DIRKQARQLE NETDJKLVSF SELCTSYHS STRGDRDRY SDDTPILNG SSQDRMFETW AIEJEQQLJAR ITCYNDKEMAE ITNSACVPSI NAALMHTLQR ERDLQDTH EFKHTKANEM AIRERENIMG SVERDIESYK SGSGSYNNRRT ELTETDIDEL RNSDRUJEET ISIAMATKEN MUSQRGMLK5 IHSKMTNLAN RUPPAVNLSIQ RINERKRD5 LILGGTIGIC TELLLLYAF3
44	SNARE	Vt11b	MGAISLTSGT QEKLIIRDDE KOEAKNMLT OMREBLHYAP VSENFNMISK LDQYQDIAQ EHLEARIMPQ DRGDMDKYGTY AVENEHMNRL QSQRAMLQG TKSLGRATQE TDQIGSEISE EIGNORDQ
45	SNARE	Membrin	MDPLFQOTHK QVHELIOSCAG RUEFTADKOSV HUYENETOAS IDQITSRIFER LETLSSKEPP NKRONAHLRV DOLKVDVQHL QPAHLRNFEHQH RHAPEQQFRQ REELLSRFET TNDSDTTPM DESLQFNSSL QRVHNGMDD1 TLDGHNILDG LTRQTLIKG TQRKJJDIAN MLGLSNIVMR LIKBRKAQDK YPMIGCMILT CYWMFLVQY LT
46	SNARE	Pallidin	MSPVGPSPSPD GALTTRPFYCI EAEQPTPGC DTSUVEGLIE DLTEDKAVF QIAEGLISHY LPDLQRSKQA LQELTNQNV LLDTLQEIS KFTECHSMID INALPAEAKH YHAKLVNIRK EMMMLHEKTS KURKRALKLQ QREKEEELR EQREKEEFER EKQLTAPAK RM
47	SNARE	Syntaxis-5	MSCRDRTQEFLSACKSLQTR QNGOTNPKA LRAYRORSEF TLMARKIGKD LSNTTAKLER HTILAKRKS1 FDDRAVEEE ITTIIKODIN SLANKOALQK DFRAKRSQS GRHQLQHNT IVVSLQSKLA SMSNDEKSVL EVERTENLKKQ RERREQESRA PVSALPAPN HLGGAIVVL AESHASHKDV4 IDMMDSRTSQ QLQILDEQD YIQSRADTMQ NIESTTVELG SIFOQLAHMV KEQEBETTQRI DENVIGAQD VEMAHSEILK YFQSYTSNWRW LMVKFLILLI VFVILEVVFL A
48	SNARE	Syntaxis-6	MSMEDDPFFV KGEVQKAINT AQGLFQRWTE LQDQSTATAR EETIWTNEJ RNNLRSTEWD LEDLDETISI VEANHKFNL DATELSIRKA FITSPRQVR DMKDQMSSTSS VOALAERKNR QNLIGDGSQ NWSTGTTDKY GRDRELQRA NSHETEEQQA OCQLTIVEQD EOLELYGSII CVIKNMQR1 CGELEEQAVM LEDESHELLES TOSRLDNVMK KLAKYSHNTS DRONCAJAI LFAYLLVYL1 IELV1
49	SNARE	Syntaxis-7	NSYTPGVGGD PAQIAQCRSS NIOKITOCSV EIGORTINOLG TPQDSPLRQ QLOOKQOQTN QIAKETDIKY1 KEGSLPTP SEDORKIKQ DRJVAEFTS LTNEFOKVRQQ AAEREKEEVA RVRASSERVSG SFPESSKER NLVSVNESQTQ PQVQVQDEE1 TEDDLRLLIE RESSIROLEA DMDINE1FK DLMGMHIEQG DVDSZEANV ENALEVHVQQA NOQLSRADY QRKSRTLCI TIIILVIGVA ITISLTWGLN H

50	SNARE	Syntaxin-8	MADPDMSTY DSTCQIAQEI AEKIQQRNQY ERGEKAPEL TUTIRALQN IKEKIALKD ILLRAVSTHQ UTQEGYRRO NLLDDLVTR EULLASTKNE GRPDPLRSQ INSREAKRGA PNPWLFEEPE ETRGLEGEDEI RQOQKILIE QDAGLDALNS LLSRQKOMGQ EIGNDLDEQN EILDOLANLV ENTEDEKLNE TRRVNMVDRX SASCGMIVYI LLLVVAIVVV AVFETN
51	SNARE	Syntaxin-10	MSLEDPPFFV RGEVQRAVNT ARGLYQRWC E LQESAAVRYI EEQOATQOLI MDEQQOLEM LEDLEETIGI VEANPGKFKL PAGDLQERKV EVERNREAVQ EMKDHMSPT AVAFIERNNR EILLAGKPAQ KSPSDLDDAS AVSATSRYE EOCATOQLIM DEODQOLEMV SGSIVNLKHM SGRVEEELDE QGIMLDAFAQ EMDHTQSRM GYLRLAKVS AMTSDRQWC ALAVIVGVLL
52	SNARE	SYNTHAXIN-10a	CAIAVNGVIL LIVLILFSL MSLEDPPFFV RGEVQRAVNT ARGLYQRWC E LQESAAVRYI EEQOATQOLI MDEQQOLEM LEDLEETIGI VEANPGKFKL PAGDLQERKV EVERNREAVQ EMKDHMSPT AVAFIERNNR EILLAGKPAQ KSPSDLDDAS AVSATSRYE EOCATOQLIM DEODQOLEMV SGSIVNLKHM SGRVEEELDE QGIMLDAFAQ EMDHTQSRM GYLRLAKVS AMTSDRQWC ALAVIVGVLL
53	SNARE	Syntaxin-11	LVLILFSL MKDRLAELLQ LSXQYDQQFP DGDDEFDSPI EDIVFETDH LESLYRDIRD IQDENIQLIVA DVKRJGKONA REFTSMRRLS SIXRUTNSLA KAVERGEVU HKCLARMKEL SEAABRQHGP HSavarisbra QYNALTIL'FQ RAMHDYNOAE MKORDNCIKR IQRQLLMGK EVSGQFLDN FEQGRKWDVFES ENLLADYKGR GPFTRSSRA TANCZAWRA IDYVHBLQ MAVLVEKQAL
54	SNARE	Syntaxin-12	TINVTELINQ KVTDYTGQK AQPRAKQYE ERNPCTLC FCCPCIK MSYGFELDMYR NGFESGPQLR DFISSLIQTS GNLRISQAT AQIKNLMSQL GTKQDQSKLQ ENLQQLQHST NQLAKETNL DSELSLPLP LSSEORQR LQERIMNDF SAALNNQFAV ORRVSEKEKE SIERRAAGRN ISSEPERORRE QLYSFSDSHE WNOQSCEDAE VARTEQDIEL IKERETAIRO LEADLIDVNO IFEDLAMMH DQGDLSLE ANVESSEVIV ERATEQLORA AYTOKKSRRK MCIIIVLVL3V LILIGLIIW LYKTK
55		Syntaxin-17	MSEDEEVKL RELPAIQKF IKVLPITNL RLRKHQINIE KYQCRJLWDK LHEEHINAGR TVQQLRSNR EZEKLCLEVR KDDIVLKKM IDPKKEEASA ATAEFLQHLI ESVELLKQF NDEETILQPP LTRSMTYGGA FHTTEAESS QSITQIYALP EIPODQNAAE SHELLEAUL ELSQUVTDFS LIVNNSQFKT DSTADHVNSA AVYVEEGTKN IGKAARYKIA ALPVAGHLIG GMVGGPIGLL ACEFKVAGIRA ALGGGYLTGET GGKLQREKKQ KMVKETLSSC PDLPSPOTDK CS
56	SNARE	VAMP-2	MSATRATAPP AAPAGEGGPP APPNPLTSNR RLQQTQAVD ENVDMRNV DKVLERDQKL SELDDRADAL QAGASQFETS AAUKRKYWW KNKEMMILLG VICALLII IVYFESS
57	SNARE	VAMP-3	MSTGPTATAVG SNRRIQQTQN QVDEVDMR VNVTVKYLERD QKLSSELDRA DALQGASQF ETSAAKLRK YWKNCMWA IGILIVVIFI LITIVWVVS

58	SNARE	VAMP-4	MPPKEFTRILN DDDVTGSIYS ERNLLEDDS JEEDEFITRG FSGPRFGPRN DKLKHVQNVQ DEVIDVMPEN ITKVIERCER LDPLQDKSES ISDNATAFSN RSKQLRQMW WRGCKRIKAIM ALVAAILLIV LILLIVMYR T
59	SNARE	VAMP-7	MALLFAVIR GTTILAKIAN CGGNFLEVTE OILAKIPSEN NKLYSHGNX LCLYLCQDRT VIJICITDDDF ERSRAFENTN EIKKREFTTY GSAQATLPY AMNSEFSVL AAGLKHSEN KGLDKYMETQ AQVDELGTM VENIDLVAQR GERLELLIDK TENLVDSVT EKTTSRNLR
60	SNARE	VAMP8	AMCMKNLKLUT LILLIVSIVT IYTJVSPICG GFTWPSCYK MEEASEGGGN DRVRNLQSEV EGVKNLNTQN VETILARGEN LEHLRNKTED LEATSEHFKT TSQKVARKFW WKNVKMIVLI CVIVFLILF IVLFATGAFS
61	SNARE	VII1-a-beta	MSDFEGIEQ DFLVTLAAT SKIARVPRLP DEEKKOMVAN VPKQLEAKE LLEQMDLEVR EIPPPQSRCM SNRMRSYKQE MCKLLETDEKR SRIAYSDEVR NELLGGDGNS SENQRHLLD INTERLERSRR RLEAGYQQLAV EUHQIGQEML BNLSHSDREKI QEARERLRET DANIGKSSRJ LTGMILRRGCS VKKQCNLSLA PKA
62	SNARE	XP350893	MRDLILPDITA CRKNDGDTV VVEKDHFMD DPFHQVEEIR NSIDKITYV EEVRENHSII LSAPAPPEFKI KEELEDLNKE IKTANKLRA KLUZALEQSFID QDESGNRTSV DLRLRBTQHS VLSRKFVAM AEYNEAQTLF RERSKGRIQR QLEITGRITT DDELEEMLES GKPSLFI'SDI ISDSQITRQA INBIESRHKD EMKLETSIRE IHEMEDMAN FVETOGEVIN NIERNVMNAT DYVEHAKETT KKAIIKYOSKA RBSVLASKN
63	SNARE	I.I.P5	CMAALAALPFP LPAQFKSICH HIFTAQEHDK RDPPVAYYCR LYAMOTGGMKI OSKTPCCKF LSKLMDOLEA LJKOLGDNEA ITQEIVGCX LENZAJKMF YADNEDAGR FHNMKSFY TASLLLIVIT VFGELTDENV KURKARYWKA TYIHNCLKEW GDSSSRPCWE LKKMILKWM KMLEQPLCPPL SQISSHHLQL MTQOHALRQL YWNTDSSGGCT RSS

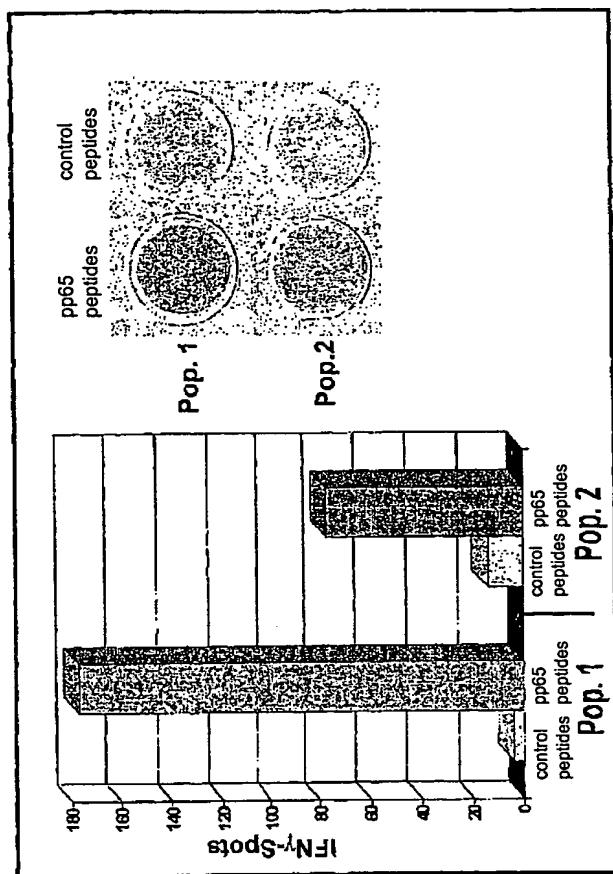


Fig. 11

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